	Gin	AST 101	.0	His	Arg	Ser	Val 101		. Val	Ile	Pro	GI u 102	_	Glu	Ala	Glu
5	Val 102		Gln	Glu	Val	Arg 103		Cys	Pro	Gly	Arg	_	Tyr	Ile	Leu	Arg
10	Val	Thr	Ala	Tyr	Lys 104		Gly	Tyr	Gly	Glu 105	_	Cys	Val	Thr	Ile 105	
	Glu	Ile	Glu	Asp 106		Thr	Asp	Glu	Leu 106	-	Phe	Ser	Asn	Cys 1070		Glu
15	Glu	Glu	Val 107		Pro	Asn	Asn	Thr		Thr	Cys	Asn	Asn 1089	_	Thr	Gly
	Thr	Gln 109	Glu 0	Glu	Tyr	Glu	Gly 109		Tyr	Thr	Ser	Arg 1100		Gln	Gly	Tyr
<b>20</b>	Asp 1105		Ala	Tyr	Gly	Asn 1110		Pro	Ser	Val	Pro 1115		Asp	Tyr	Ala	Ser 1120
1 25	Val	Tyr	Glu	Glu	Lys 1125		Tyr	Thr	Asp	Gly 1130	_	Arg	Glu	Asn	Pro 1135	_
	Glu	Ser	Asn	Arg 1140		Tyr	Gly	Asp	Tyr 1145		Pro	Leu	Pro	Ala 1150		Tyr
30	Val	Thr	Lys 1155		Leu	Glu	Tyr	Phe 1160		Glu	Thr	Asp	Lys 1165		Trp	Ile
	Glu	Ile 1170	Gly	Glu	Thr		Gly 1175		Phe	Ile		Asp 1180		Val	Glu	Leu
35	Leu 1185		Met	Glu	Glu											
40	(2)	INFO	RMAT	lon	FOR	SEQ	ID N	IO : 5 :								
		(i)		L) LE	ENGTH	IARAC I: 35 nucl	67 b	ase	pair	s						
45						EDNE GY:		_	rle							
		(ix)		) NA	ME/F	ŒY:		3567								
50		(xi)	SEC	UENC	E DE	ESCRI	PTIC	N: S	EQ I	D NC	):5:					

	Glu		Gln					AGT Ser		48
5								AAT Asn		96
10	TCA Ser								1	144
15	GTA Val 50								1	192
20	ATA Ile								2	240
Company   Comp	TTA Leu								2	88
25	AAT Asn								3	36
TO THE THE PARTY OF THE PARTY O	AAA Lys								3	84
	ATT Ile 130									32
40	TCG Ser								4	80
. •	CAA Gln	-							5	28
45	GGA Gly								5	76
50	AGA Arg								6	24

		Asn						Thr			GAT Asp	672
5	Ile	ACA Thr									TTA Leu 240	720
10		GCC Ala										768
15		GTT Val										816
20		AAT Asn 275										864
20 10 10 25 25		GAG Glu										912
25		CTT Leu										960
3 <b>6</b>		GGA Gly						_	_	_		1008
35		TCT Ser										1056
40		ACT Thr 355										1104
40		TTA Leu	_									1152
45	 	GAA Glu		 	 							1200
50		AGA Arg										1248